1/4

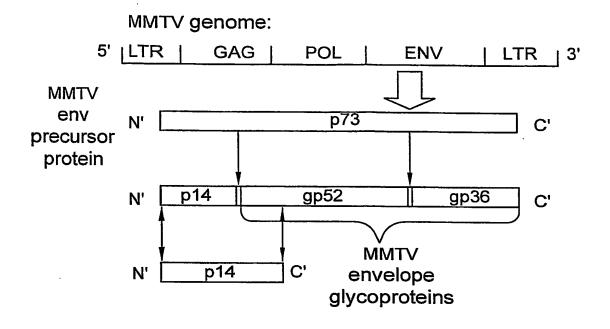


FIG. 1A

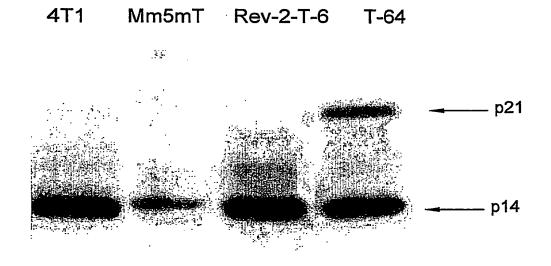
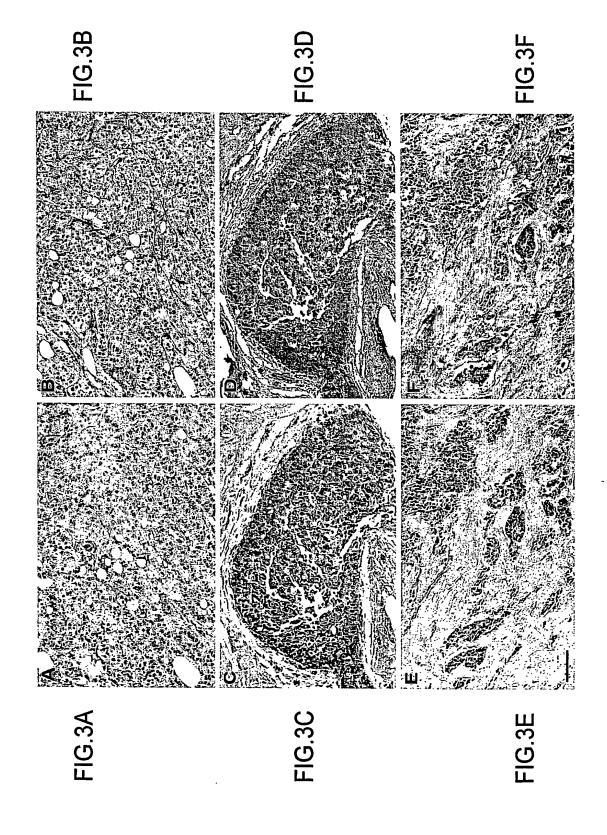


FIG. 1B

7 GSPTGSSDLLLSGKKQRPHLALRRKRRREMKKINRKVRRMNLDLIKEKTA 56 2 GSLTDSSDLSLSGKDRRPRLALRRKRRREMRKITKKVRRMTLDPIKEKTA 93 52 WQQLQALISEAEEVLKTSQTPQTSLTLFLALLSVLGPLPVTG 57 WQHLQALISEAEEVLKTSQTPQTSLTLFLALLSVLGPPVTG af248270 p14

FIG. 2



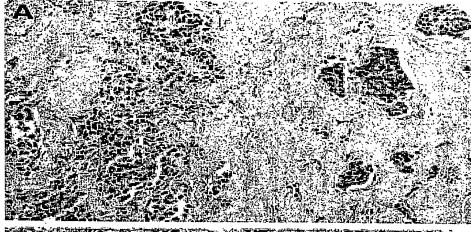


FIG. 4A

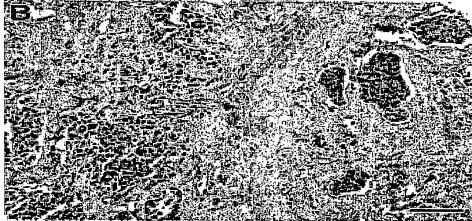


FIG. 4B

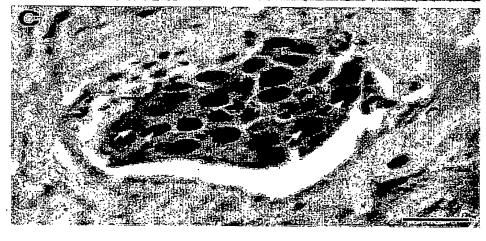


FIG. 4C